

| Result | No. | Score | Query | Match | Length | DB | ID | Description |
|--------|-----|-------|-------|-------|--------|--------|----|-------------------|
| C | 1 | 708 | 98.6 | 718 | 21 | A05545 | | Human Immunogeni |
| | 2 | 513.2 | 71.5 | 812 | 21 | A06690 | | Human Immunogeni |
| | 3 | 513.2 | 71.5 | 820 | 19 | V62429 | | Prostate cancer a |
| | 4 | 513.2 | 71.5 | 1872 | 19 | V62428 | | Prostate cancer a |
| | 5 | 457.6 | 63.7 | 3112 | 21 | A06687 | | Human immunogeni |
| | 6 | 457.4 | 63.7 | 2229 | 21 | A06688 | | Human immunogeni |
| | 7 | 457.4 | 63.7 | 2426 | 21 | A06689 | | Human immunogeni |
| | 8 | 457.2 | 63.7 | 2037 | 19 | V62427 | | Prostate cancer a |
| | 9 | 457.2 | 63.7 | 3582 | 19 | V62430 | | Prostate cancer a |
| | 10 | 289.8 | 40.4 | 359 | 20 | X37445 | | Human prostate c |
| | 11 | 257 | 35.8 | 597 | 20 | X37486 | | Human secreted p |
| | 12 | 93.8 | 13.1 | 301 | 21 | A05520 | | Human immunogeni |

[illegible]

CC was found in approximately 15% of the cDNA clones isolated. The
CC invention claims for PCa3 CDNA variants and the proteins they encode.
CC The invention also claims for antibodies against PCa3 protein. The
CC antibodies are claimed to be useful for detecting PCa3 protein in
CC immunassay tests, for diagnosing, assessing and prognosing of
CC prostatic cancer (PC). Antibodies, optionally coupled to a cytotoxin
CC or radioisotope, and nucleic acids antisense to PCa3 cDNA are claimed
CC to be useful for treating PC, while determining elevated levels of
CC PCa3 (as RNA or protein) is useful for detecting a predisposition
CC to development of PC, e.g. in prenatal tests. Detecting PCa3 protein
CC allows differentiation between malignant and benign prostatic disease,
CC and the level of PCa3 expression allows correlation with the grade of
CC tumour. PCa3 protein and its fragments are also claimed to be useful
CC in vaccines for preventing PC; in drug screens for identifying
CC specific (ant)agonists (potentially useful therapeutically) and for
CC studying protein-DNA interactions.

XX Sequence 820 BP; 262 A; 169 C; 191 G; 198 T; 0 other;

XX

Query Match 71.5%; Score 513.2; DB 19; Length 820;
Best Local Similarity 97.3%; Pred. No. 2,3e-148;
Matches 585; Conservative 0; Mismatches 9; Indels 7; Gaps

OY 1 ggaagattgtgtgtgttcgcagccggaggagaccaagaaatctgcattggtggaagacc 60
DB 27 ggaagattgtgtgtg-ctgcagcgcggggagacacagaaatctgcattgtggaagacc 85
OY 61 tgaataacacagaggtggaataataagaagaagcgcgcgcatttaccatcctgaagccacaact 120
DB 86 tgaatatacagaagtggaataataagaagaagcgcgcgcatttaccatcctgaagccacaact 145
OY 121 ctgtctgaataggagaataataacatcactatagaacaagcaagaatgacaatataatgtctaa 180

| ID | RESULT | standard; cDNA; 1872 bp. | |
|----|--------|---|-----|
| QY | 181 | gtatgtgacatcgttttttgcacatcttccagccctttaaatatccacacacacagagaagac | 240 |
| Db | 206 | gtatgtgacatcgttttttgcacatcttccagccctttaaatatccacacacagagaagac | 265 |
| QY | 241 | aaagagaagcacaagagatccctctggagaaatgcccggccgcacattcttggtcatcgatga | 300 |
| Db | 266 | aaagagaagcacaagatccctctggagaaatgcccggccgcacattcttggtcatcgatga | 325 |
| QY | 301 | gctctgcacctgtgcctgnttcccgctgtgtgagggaaaggagacattagaagaaatgaaatgtgtg | 360 |
| Db | 326 | gctctgcacctgtgcctgnttcccgctgtgtgagggaaaggagacattagaagaaatgaaatgtgtg | 385 |
| QY | 361 | ttcccttaagagat-ggcagagaaacagatcccgctgtgtgatacttaatttgaaacggagatta | 419 |
| Db | 386 | ttcccttaagagatgaggcagagaaacagatcccgctgtgtgatacttaatttgaaacggagatta | 445 |
| QY | 420 | cagatttgaaaatgaaagtcacaaagatgaaagcatctaccaaataagaggaagaaacagagaaaaa | 479 |
| Db | 446 | cagatttgaaaatgaaagtcacaaagatgaaagcatctaccaaataagagggaaaacagagaaaaa | 505 |
| QY | 480 | tcttgatgtg-ttccacaagacatgcaacaacaacaaatgtgatactgtgtgaatgacatgagga | 536 |
| Db | 506 | tcttgatgtgtccacacaagacatgcaacaacaacaaatgtgatactgtgtgaatgacatgagga | 565 |
| QY | 537 | agccaacatcgggagagagat-accacggggcaga-gtccaagatctcttgccctgtgccta | 594 |
| Db | 566 | agccaagctgggagagagataaacacacggggcagaggtgtccaagatctcttgccctgtgccta | 625 |
| QY | 595 | a 595 | |
| Db | 626 | a 626 | |

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AC V62428;
XX
XX 30-DEC-1998 (first entry)
DE Prostate cancer antigen (PCA3) cDNA splice variant 2.
XX
XX Prostate cancer antigen cDNA splice variant 2; PCA3; prostatic cancer;
XX PC; ds.
XX
XX Homo sapiens.
XX
XX WO9845420-A1.
XX
XX 15-OCT-1998.
XX
XX 09-APR-1998; 98WO-CA00346.
XX
XX 10-APR-1997; 97US-0041836.
XX
XX (DIAG-) DIAGNOCURE INC.
XX
XX Bussemakers MFG;
XX
XX WPI; 1998-568347/48.
XX
XX New nucleic acid encoding prostate cancer antigen 3 - for diagnosis,
XX prevention and treatment of prostatic cancer
XX
XX Claim 4; Pages 76-77; 11pp; English.
XX
XX The present sequence represents the prostate cancer antigen (PCA3)
XX cDNA splice variant 2 sequence comprising of exons 1, 3, 4a and
XX 4b of the PCA3 gene. The PCA3 cDNA splice variant 2 sequence,
XX isolated from a human primary prostatic tumour tissue cDNA library,
XX was found in approximately 65% of the cDNA clones isolated. The
XX invention claims for PCA3 cDNA variants and the proteins they encode.
XX The invention also claims for antibodies against PCA3 protein. The
XX antibodies are claimed to be useful for detecting PCA3 protein in
XX immunosay tests, for diagnosing, assessing and prognosing of
XX prostatic cancer (PC). Antibodies, optionally coupled to a cytotoxin
XX or radioisotope, and nucleic acids antisense to PCA3 cDNA are claimed
XX to be useful for treating PC, while determining elevated levels of
XX PCA3 (as RNA or protein) is useful for detecting a predisposition
XX to development of PC, e.g. in prenatal tests. Detecting PCA3 protein
XX allows differentiation between malignant and benign prostatic disease.
XX and the level of PCA3 expression allows correlation with the grade of
XX tumour. PCA3 protein and its fragments are also claimed to be useful
XX in vaccines for preventing PC; in drug screens for identifying
XX specific (ant)agonists (potentially useful therapeutically) and for
XX studying protein-DNA interactions.
XX
XX Sequence 1872 BP; 567 A; 389 C; 369 G; 539 T; 8 other;
SQ

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Query Match 71.5%; Score 513.2; DB 19; Length 1872;
 Best Local Similarity 97.3%; Pred. No. 3.6e-148;
 Matches 585; Conservative 0; Mismatches 9; Indels 7; Gaps 6;

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OY 1 ggaattgtgtgttgcagccgaggaagcaggaagatctcagtggtgggaagacc 60
   |||||||
Db 27 ggaattgtgtgtg-cgcgacccgaggaagcaggaagatctcagtggtgggaagacc 85
OY 61 tgatgatacagaggtggaagaataagaagagctgtgacttaccatcttgaggccacacat 120
   |||||||
Db 86 tgatgatacagaggtggaagaataagaagagctgtgacttaccatcttgaggccacacat 145
OY 121 ctctgtgaatgagagataataacatacactagaacacagacagatgacaataatgtctaa 180
   |||||||
Db 146 ctgtgaatgagagataataacatacactagaacacagacagatgacaataatgtctaa 205
OY 181 gtatgtacatgttttgcacattccagccctttaataatccacacacaggaagacac 240
   |||||||
Db 206 gtatgtacatgttttgcacattccagccctttaataatccacacacaggaagacac 265

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OY 241 aaaagaagcacagagatcccttgaggagaatgccgcgccatcttggtcatcatgta 300
   |||||||
Db 266 aaaagaagcacagagatcccttgaggagaatgccgcgccatcttggtcatcatgta 325
OY 301 gccctgcctgtgctgtcccgctgtgagggaagagacattagaataatgattgattg 360
   |||||||
Db 326 gccctgcctgtgctgtcccgctgtgagggaagagacattagaataatgattgattg 385
OY 361 ttcccttaaggat-ggcaggaanaacagatccctgtgtgtgatatattatttgacaggat 419
   |||||||
Db 386 ttcccttaaggatgagcaggaanaacagatccctgtgtgtgatatattatttgacaggat 445
OY 420 cagatttgaatgaatgacatacaagaatgacattaccatgtagaggaagaaacagagaaa 479
   |||||||
Db 446 cagatttgaatgaatgacatacaagaatgacattaccatgtagaggaagaaacagagaaa 505
OY 480 tcttgatgg-ttcaacaagatgcacaacaagaatgatactgtgtagacagag--c 536
   |||||||
Db 506 tcttgatggcttcaacaagatgcacaacaagaatgatactgtgtagacagagca 565
OY 537 agccaactggggaagagat-accacggggcaga-gtcaagattctggccctgtgctta 594
   |||||||
Db 566 gccaaactggggaagagataaccacggggcagagaggtcagattctggccctgtgctta 625
OY 595 a 595
Db 626 a 626

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RESULT 5
 A06687
 ID A06687 standard; cDNA; 3112 BP.
 AC A06687;
 XX
 DT 13-JUN-2000 (first entry)
 XX
 DE Human immunogenic prostate tumour protein cDNA sequence SEQ ID NO:468.
 XX
 XX Human; prostate cancer; diagnosis; tumour; gene therapy; detection;
 KW Immunogenic; cyostatic; vaccine; ss.
 OS Homo sapiens.
 PN WO200004149-A2.
 XX
 XX 27-JAN-2000.
 PD
 XX
 XX 14-JUL-1999; 99WO-US15838.
 PF
 XX
 PR 14-JUL-1998; 98US-0115453.
 PR 14-JUL-1998; 98US-0116134.
 PR 23-SEP-1998; 98US-0159812.
 PR 23-SEP-1998; 98US-0159822.
 PR 15-JAN-1999; 99US-0232149.
 PR 15-JAN-1999; 99US-0232880.
 PR 09-APR-1999; 99US-0288946.
 XX
 XX (CORI-) CORIXA CORP.
 XX
 XX Dillon DC, Harlocker SL, Yugu J, Xu J, Mitcham JL;
 PI
 XX
 DR WPI; 2000-171268/15.
 XX
 PT New polypeptide useful for treating and diagnosing prostate cancer
 PT comprises an immunogenic portion of prostate tumor protein -
 XX
 PS Claim 1; Page 259-260; 263pp; English.
 XX
 CC The present invention describes isolated polypeptides, comprising an
 CC immunogenic portion of a prostate tumour protein (PTP). The polypeptides
 CC and polynucleotides encoding them have cyostatic activity and can be

Sequence 3112 BP; 975 A; 587 C; 624 G; 926 T; 0 other;

67 taacagagtgaataagaagcgtctgacttaccatctgagccacacctgctg 126
||| ||||||||||||||||||||||||||||||||||||||||||||
1307 taataagctgagaataagaagcgtctgacttaccatctgaggcacacacctgctg 1366

| | |
|----------|---------------------------------|
| RESULT | 6 |
| A06688/c | |
| ID | A06688 standard; cDNA; 2229 BP. |

| | |
|----|--|
| AC | A06688; |
| XX | |
| DT | 13-JUN-2000 (first entry) |
| XX | |
| DE | Human immunogenic prostate tumour protein cDNA sequence SEQ ID NO:469. |
| XX | |
| KW | Human; prostate cancer; diagnosis; tumour; gene therapy; detection; |
| KW | immunogenic; cytostatic; vaccine; ss. |
| XX | |
| OS | Homo sapiens. |
| XX | |
| PN | WO200004149-A2. |
| XX | |
| PD | 27-JAN-2000. |

| | |
|----|---|
| PI | Dillon DC, Harlocker SL, Yuguu J, Xu J, Mitcham JL, |
| XX | |
| DR | WPI; 2000-171268/15. |

| | | | | | | | |
|-----------------------|-------|--------------|--------------|------------|----|--------|------|
| Query Match | 63.7% | Score | 457.4 | DB | 21 | Length | 2229 |
| Best Local Similarity | 94.1% | Pred. | No. 6.1e-131 | | | | |
| Matches | 529 | Conservative | 0 | Mismatches | 27 | Indels | 6 |
| | | | | | | Gaps | 5 |

QY 40 atctgcagtggtgggaagaagacttgatgatacagaagtgaaagaataagaagaagtcgcgcgact 99

Db 1808 ATTTCCTCCCAATATGGTCTGTATGTATTTCACAGTGAAGAATAATGAAGAGCGCTGTACT 174

OY 100 ttacacatctggagccacacatctgcctaagaatggagatgaatataacataacataagaagaacga 159

Db 1748 TTACCATCTTGAGGCCACACATCTCTGTGAATGGAGATTAATTAACATCTAGTAAGAACGCA 168

OY 160 agatgacataataatgcttaagtagtgacatgcttcttgacaatctccagcccttcaat 219

Db 1668 AGATACAAATTAATGCTGTAAGTAGTACATGTTTGGCATTTTCAGGCCCTTTAAAT 162

OY 220 atccacacacacaggaagcacacaaaggaaagacacagagatctcccttggagaatctgccgcgc 279

Db 1628 ATCCACACACACAGGAAGCACAAAGGAAGACAGAGATCCCTGGAGAAATCCCGGCC 156

OY 260 gccatcttgggtcatalcgatgtagcctcgcccttgctcgtatccgctgtgtgaggaaagac 339

Db 1568 GCCATCTTGGGTCAATCGATGAGCGCTCCGCCCTGCTCCGTGCCGCTGTGAGGAAGAGAC 150

OY 340 attgagaatgaattgatatgcttcccttaagaagat-ggcagagaanaacagatcccgtctgtg 398

Db 1508 ATTGAAAAATGAATTAATGATGCTTCTTTAAAGGAATGGGACAGAAACAGATCTGTTGTGTG 144

OY 399 atattatcttaacggggtatcacatcttgaatgaagtcacaagaatgagcatctaacatcg 458

Db 1448 ATATTATTATTAAACGGGATTTCACATTTGAAATGAAGTCAAAAGTGAAGCATTAACAAATG 138

OY 459 agaggaanaacagacagagaaaaatctctga-tgg-ttcacagaagcatgcacacaaacaaaatcgga 517

Dd 1388 AGAGAAAAACGACGAAGAAATCTTGATGGGTCTTCAACAACATGCATAAACCMAAATTGTA 1329

Oy 518 atactcgtatgacacag--cacccaactggaggagaat-accacygggcaga-gtca 573
Dd 1338 ATACTGTATGTCACATGAGCAGCCAACGCTGGGAGAGAATCAACCGGGCACAGAGGTCA 1269

Oy 574 gattcttgacctgtcgcccta 595
Dd 1268 GGATTCTGGCCCCCTGCTGCCTTAA 1247

RESULT - 7
ID A06689/c
XX A06689 standard; cDNA; 2426 BP.
XX A06689;
X AC
XX AD
DT 13-JUN-2000 (first entry)
XX DE
DE Human immunogenic prostate tumour protein cdna sequence SEQ ID NO:470.

K Humam; prostate cancer; diagnosis; tumour; gene therapy; detection;
K immunogenic; cytosolic; vaccine; ss.
OS Homo sapiens.
PN WO200004149-A2.
XX PD
PD 27-JAN-2000.
XX PF
PF 14-JUL-1999; 99WO-US15838.
XX PR
PR 14-JUL-1998; 98US-0115453.
PR 14-JUL-1998; 98US-0116134.
PR 23-SEP-1998; 98US-0159812.
PR 23-SEP-1998; 98US-0159822.
PR 15-JAN-1999; 98US-0232149.
PR 15-JAN-1999; 98US-0232860.
PR 09-APR-1999; 99US-0288946.

(CORI-) CORIXA CORP.
PA Dillion DC, Harlocker SL, Yugu J, Xu J, Mitcham JL;
PI WPI; 2000-171268/15.
PT New polypeptide useful for treating and diagnosing prostate cancer
PT comprises an immunogenic portion of prostate tumor protein -
F Claim 1; Page 261-262; 263pp; English.

The present invention describes isolated polipeptides, comprising an immunogenic portion of a prostate tumour protein (PP), The polipeptides and poly nucleotides encoding them have cytostatic activity and can be used in vaccines and in gene therapy. The polipeptides and poly nucleotides encoding them, antigen presenting cells which express the polipeptides, antibodies against the polipeptides and vaccines comprising them can be used for inhibiting the development of prostate cancer in a patient. The polipeptides can be used to generate antibodies or anti-idiotypic antibodies for passive immuno therapy. A portion of the polynucleotides encoding the polipeptides can be used as a probe or reagent to modulate the expression of the polipeptides. A06241 to A06691 and CC present invention.

Sequence 2426 BP; 717 A; 476 C; 548 G; 685 T; 0 other;

| Query Match | 63.7% | Score 457.4 | DB 21 | Length 2426 |
|---------------------------|-------|--------------------|----------|-------------|
| Best Local Similarity | 94.1% | Pred. No. 6.3e-131 | | |
| Matches 529; Conservative | 0 | Mismatches 27 | Indels 6 | Gaps 5 |

0Y 40 atctgcattgttgtaagagacctgatgatcaacaggttgagaataataagaagatcgtccgaact 99

| | | | |
|----|------|--|------|
| Db | 1802 | ATTTCCTCCCAATAGTGCTGATGTATTTCCAGTGAGAAATTAAGAAAGCGTCGACT | 1743 |
| QY | 100 | ttaccatcttgaggccacacacatctctcgtgaatggaataataacatcactagaacaagca | 159 |
| Db | 1742 | TTACACTCTGGGGCCACACACACTCTGCTTAATTTAAGAGATTAATTAACTACCTACGAACACACA | 1683 |
| QY | 160 | agatatcaataaattgtcctaagtatgacatgttttttcgaatctccagcccttttaat | 219 |
| Db | 1682 | AGATACACAATTAATGTCTAAGTAGTAGACATGTTTTTGCACATTTCCAGCCCTTTAAT | 1623 |
| QY | 220 | atccacacacacaggaagcacaaagaagcacagaagatcccttggaggaatgcccggcc | 279 |
| Db | 1622 | ATCCACACACACAGAGAGCACAAAGAAGAACACAGAGATCCCTGGGAGAAATGCCGGCC | 1563 |
| QY | 280 | gccactctgggtactgatgtagcctcgccctgctgacctgttcccgcttggaggaagac | 339 |
| Db | 1562 | GCCACTCTGGGTACTGATGATAGATAGCGCTCGCCCTGTGCTGCCGTGCCGTTGAGAGGAGAGAC | 1503 |
| QY | 340 | attagaanaatgaatltgatgtgtcccttaaaagat-gccaggaanaacagatccctgtgttgg | 398 |
| Db | 1502 | ATTGGAANAATGAATTGATGTGTCTCTTAAGATGGGCGAGAAAACAGATCTCTGTGGG | 1443 |
| QY | 399 | atattatttgaacgggattacagatttgaatgaagtacaaatgacatltaccaatg | 458 |
| Db | 1442 | ATATTATTTCCTAACGGGATTACAGATTTGAATGAAGTACAAATGAGCATTTACCAATG | 1383 |
| QY | 459 | agaggaanaacagagagaanaatcttgaatg-ttcaaaagacatgaaanaacaaatgga | 517 |
| Db | 1382 | AGAGGAAAACAGACGAGAAAATTCTTAGTGCTTCACAAACATGCAACCAAACTAAATGGA | 1323 |
| QY | 518 | atactgtgatgacagcg--cagccaacttgggagagagat-accacggggcaga-gtcca | 573 |
| Db | 1332 | ATACTGTGATGACATGAGGACGACCAAGCTGGGAGAGATATCAACGCGGCAGAGGGTCA | 1263 |
| QY | 574 | ggattctggccctgctgtgacctaa | 595 |
| Db | 1262 | GGATTCTGGCCCTGCTGTCCCTAA | 1241 |

| RESULT | ID | standard | CDNA | BP. |
|--------|--|------------------------------------|------|----------|
| V62427 | V62427 | standard | CDNA | 2037 BP. |
| XX | V62427; | | | |
| XX | 30-DEC-1998 | (first entry) | | |
| XX | Prostate cancer antigen (PCA3) | CDNA splice variant 1. | | |
| XX | Prostate cancer antigen CDNA splice variant 1; | PCA3; prostatic cancer | | |
| XX | PC; ds. | | | |
| XX | Homo sapiens. | | | |
| XX | Key | Location/Qualifiers | | |
| XX | CDS | 379..534 | | |
| XX | FT | /*tag= a | | |
| XX | FT | /product= "PCA3 protein variant 1" | | |
| XX | FT | 2019..2024 | | |
| XX | polya_signal | /tag= b | | |
| XX | MO9845420-A1. | | | |
| XX | 15-OCT-1998. | | | |
| XX | 09-APR-1998; | 98MO-CA00346. | | |
| XX | 10-APR-1997; | 97US-0041836. | | |
| XX | (DIAG-) | DIAGNOCURE INC. | | |
| XX | Bussemakers MJG; | | | |
| XX | | | | |

XX MPI: 1998-568347/48.
DR P-PSDB; W79736.

XX New nucleic acid encoding prostate cancer antigen 3 - for diagnosis,
PT prevention and treatment of prostatic cancer

XX Claim 3: Fig 2B-2J; 111pp; English.

XX The present sequence represents the prostate cancer antigen (PCa3)
CC cDNA splice variant 1 sequence comprising of exons 1, 2, 3, 4a and
CC 4b of the PCa3 gene. The PCa3 cDNA splice variant 1 sequence,
CC isolated from a human primary prostatic tumour tissue cDNA library,
CC was found in approximately 5% of the cDNA clones isolated. The
CC invention claims for PCa3 cDNA variants and the proteins they encode.
CC The invention also claims for antibodies against PCa3 protein. The
CC antibodies are claimed to be useful for detecting PCa3 protein in
CC immunoassay tests, for diagnosing, assessing and prognosing of
CC prostatic cancer (PC). Antibodies, optionally coupled to a cytotoxin
CC or radioisotope, and nucleic acids antisense to PCa3 cDNA are claimed
CC to be useful for treating PC, while determining elevated levels of
CC PCa3 (as RNA or protein) is useful for detecting a predisposition
CC to development of PC, e.g. in prenatal tests. Detecting PCa3 protein
CC allows differentiation between malignant and benign prostatic disease,
CC and the level of PCa3 expression allows correlation with the grade of
CC tumour. PCa3 protein and its fragments are also claimed to be useful
CC in vaccines for preventing PC; in drug screens for identifying
CC specific (ant)agonists (potentially useful therapeutically) and for
CC studying protein-DNA interactions.

SO Sequence 2037 BP; 622 A; 426 C; 406 G; 575 T; 8 other;

Query Match 63.7%; Score 457.2; DB 19; Length 2037;
Best Local Similarity 97.2%; Pred. No. 6.7e-131;
Matches 518; Conservative 0; Mismatches 9; Indels 6; Gaps 5;

QY 69 cagaggtggaataagaaggtgctgacttaccatctgagcgacaacatctgtcga 128
DB 259 caggggtggaataagaaggtgctgacttaccatctgagcgacaacatctgtcga 318
QY 129 atggaataataatacctactagaacagcaaatgacataatgactaagtatgac 188
DB 319 atggagataataatacctactagaacagcaaatgacataatgactaagtatgac 378
QY 189 atgttttgcacattccagccctttaataatccacacacaggaagagcaaaagaa 248
DB 379 atgttttgcacattccagccctttaataatccacacacaggaagagcaaaagaa 438
QY 249 gcaagaagaatccctgggagaatgcccggccgacatctgggtcatcgatgagctgcg 308
DB 439 gcaagaagaatccctgggagaatgcccggccgacatctgggtcatcgatgagctgcg 498
QY 309 ctgtgacctgtccgcttctgaggaagacattagaataatgattgattgttctttaa 368
DB 499 ctgtgacctgtccgcttctgaggaagacattagaataatgattgattgttctttaa 568
QY 369 aggat-ggaggaagaaacagatcctgttggatattatttgaacggattacagattg 427
DB 559 aggatggaggaagaaacagatcctgttggatattatttgaacggattacagattg 618
QY 428 aaatgaagtcacaaagtgtgacattaccatgagaggaagaaacagaaatcttgatg 487
DB 619 aaatgaagtcacaaagtgtgacattaccatgagaggaagaaacagaaatcttgatg 678
QY 488 g-ttcacagacatgcaacaacaataatggaatctgtatgacagag--cagccaact 544
DB 679 gcttccacagacatgcaacaacaataatggaatctgtatgacagagcagcagcagct 738

QY 545 gggagagagat-accacggggcaga-ggtcagagatcttgccctgtgcttaa 595
DB 739 gggagagagataaccacggggcagaggtcagagatcttgccctgtgcttaa 791

RESULT 9
ID V62430 standard; CDNA; 3582 BP.
XX V62430;
AC V62430;
XX

DT 30-DEC-1998 (first entry)

XX Prostate cancer antigen (PCa3) wild-type cDNA.

XX Prostate cancer antigen cDNA; PCa3; prostatic cancer;
KM PC; ds.

XX Homo sapiens.

XX Key Location/Qualifiers
FH CDS 401..556
FT /tag= a
FT /product= "PCa3 protein"

FT polyA_signal
FT 983..987
FT /tag= b

FT polyA_signal
FT 2041..2046
FT /tag= c

FT polyA_signal
FT 2597..2602
FT /tag= d

FT polyA_signal
FT 3494..3496
FT /tag= e

XX MO9845420-A1.

XX 15-OCT-1998.

XX 09-APR-1998; 98MO-CA00346.

XX 10-APR-1997; 97US-0041836.

XX (DIAG-) DIAGNOCURE INC.

XX Bussemakers MUG;

XX MPI: 1998-568347/48.

XX P-PSDB; W79736.

XX New nucleic acid encoding prostate cancer antigen 3 - for diagnosis,
PT prevention and treatment of prostatic cancer

XX Claim 3: Fig 5B-5F; 111pp; English.

XX The present sequence represents the prostate cancer antigen (PCa3)
CC wild-type cDNA sequence comprising of exons 1, 2, 3, 4a-4d of the
CC PCa3 gene. The invention claims for PCa3 cDNA variants and the
CC proteins they encode. The invention also claims for antibodies
CC against PCa3 protein. The antibodies are claimed to be useful for
CC detecting PCa3 protein in immunoassay tests, for diagnosing, assessing
CC and prognosing of prostatic cancer (PC). Antibodies, optionally
CC coupled to a cytotoxin or radioisotope, and nucleic acids antisense
CC to PCa3 cDNA are claimed to be useful for treating PC, while determining
CC elevated levels of PCa3 (as RNA or protein) is useful for detecting a
CC predisposition to development of PC, e.g. in prenatal tests. Detecting
CC PCa3 protein allows differentiation between malignant and benign
CC prostatic disease, and the level of PCa3 expression allows correlation
CC with the grade of tumour. PCa3 protein and its fragments are also
CC claimed to be useful in vaccines for preventing PC; in drug screens
CC for identifying specific (ant)agonists (potentially useful
CC therapeutically) and for studying protein-DNA interactions.

SO Sequence 3582 BP; 1052 A; 788 C; 679 G; 1063 T; 0 other;

Query Match 63.7%; Score 457.2; DB 19; Length 3582;
Best Local Similarity 97.2%; Pred. No. 8.9e-131;
Matches 518; Conservative 0; Mismatches 9; Indels 6; Gaps 5;

| | | | |
|-----------|--|---|-----|
| Qy | 69 | cagaggttagaataagaagaagcgctgacttaccctctcgtaggccacaactctgtgaa | 128 |
| Db | 281 | caaggggttgagaaataagaagcgctgacttaccctctcgtaggccacaactctgtgaa | 340 |
| Qy | 129 | atgagagataataacatcactagaaacagacagaatgacaataatctcaatagtgc | 188 |
| Db | 341 | atggagataataatacatcactagaaacagacagaatgacaataatctcaatagtgc | 400 |
| Qy | 189 | atgtttttgcaatttccagccctttaataatccacacacacaggaagcacaaaagaa | 248 |
| Db | 401 | atgtttttgcaatttccagccctttaataatccacacacacaggaagcacaaaagaa | 460 |
| Qy | 249 | gcacagagatccctctggagaaatcccgccgcacattgttgcatcagatgagcctgcgc | 308 |
| Db | 461 | gcacagagatccctctggagaaatcccgccgcacattgttgcatcagatgagcctgcgc | 520 |
| Qy | 309 | ctgtgctgntcccgctgtgtgagggaagacatagaaaaatgatgatgtgtccttaa | 368 |
| Db | 521 | ctgtgctgntcccgctgtgtgagggaagacatagaaaaatgatgatgtgtccttaa | 580 |
| Qy | 369 | aggat-ggcacggaagaacacagatccctgtgtgataattatgacgggattacagattg | 427 |
| Db | 581 | aggatgggcaggaagaacacagatccctgtgtgataattatgacgggattacagattg | 640 |
| Qy | 428 | aaatgaatccaaagttagcattaccatagagagaaacagacagaagaatcttgatg | 487 |
| Db | 641 | aaatgaatccaaagttagcattaccatagagagaaacagacagaagaatcttgatg | 700 |
| Qy | 488 | g-ttacaagaacatgcacaacaacaaatggatactgtgtgacagag--cagcccaact | 544 |
| Db | 701 | gcttacaagaacatgcacaacaacaaatggatactgtgtgacagagcagcgt | 760 |
| Qy | 545 | ggggagggagat-accacggggcaca-ggtcaagattctggcctctgcctaa | 595 |
| Db | 761 | ggggagggagataccacacggggcagaggtcagatctggcctctgcctaa | 813 |
| RESULT 10 | | | |
| ID | 233445 | standard; cDNA; 359 BP. | |
| XX | 233445; | | |
| AC | 233445; | | |
| XX | | | |
| DT | 08-DEC-1999 | (first entry) | |
| XX | | | |
| DE | Human prostate cancer-associated EST 23. | | |
| XX | | | |
| KX | Expressed sequence tag; EST; prostate tumor; antitumor; treatment; | | |
| KX | gene therapy; tissue specificity human; ss. | | |
| XX | | | |
| OS | Homo sapiens. | | |
| XX | | | |
| PN | DE19811193-A1. | | |
| PD | 16-SEP-1999. | | |
| XX | | | |
| PF | 10-MAR-1998; 98DE-1011193. | | |
| XX | | | |
| PR | 10-MAR-1998; 98DE-1011193. | | |
| XX | | | |
| PA | (META-) METAGEN GES GENOMFORSCHUNG MBH. | | |
| PI | Specht T, Hinzmann B, Schmitt A, Pilarsky C, Dahl E, Rosenthal A; | | |
| DR | WPI: 1999-519628/44. | | |
| XX | | | |
| DR | P-PSDB; Y48243. | | |
| XX | | | |
| PT | New nucleic acid expressed at high level in prostatic tumor tissue and | | |
| PT | encoded polypeptides, useful for treating cancer and screening for | | |
| PT | therapeutic agents | | |
| XX | | | |
| XX | Claim 1a; 87; 16pp; German. | | |

| | |
|--|--|
| XX | This invention describes novel nucleic acid sequences (A) that are |
| CC | expressed at high level in prostatic tumor tissue and encode gene |
| CC | products or their fragments. The products of the invention have |
| CC | antitumor activity. Polypeptides (I) encoded by (A) are used: (i) for |
| CC | identifying agents for treatment of prostatic cancer and (ii) for therap |
| CC | of prostate cancer, optionally where expressed by gene therapy methods. |
| CC | (A) is also used to isolate full-length genes (for gene therapy) and |
| CC | for recombinant production of (i), which can be used to raise specific |
| CC | antibodies. (A) are identified by assembly of ESTs (expressed sequence |
| CC | tags) before they are analyzed for expression pattern (tissue |
| CC | specificity). This approach eliminates many of the false results, as |
| CC | regards tissue specificity, associated with known methods that use |
| CC | single (usually short) ESTs. Z33423-Z33476 represent expressed sequence |
| CC | tags described in the method of the invention. |
| XX | |
| SQ | Sequence 359 BP; 121 A; 75 C; 94 G; 69 T; 0 other; |
| | |
| Query Match | 40.4%; Score 289, 8; DB 20; Length 359; |
| Best Local Similarity | 99.0%; Pred. No. 1e-79; |
| Matches 302; Conservative 0; Mismatches 2; Indels 1; Gaps | |
| OY | 1 ggaagattgttgttgcttcacgcgaggagagacagaagatctgatgtggaaagacc 60 |
| Db | 56 ggagattctgtgtg-ctcgacgcgaggagagaccagaagatctcatgtggaagagacc 114 |
| OY | 61 tgatatcacagagtgtgaanaataagaagcgtgtgtgactttaccatcttagggccacacat 120 |
| Db | 115 tgaigtatcacagagtgtgaanaataagaagcgtgtgtgactttaccatcttagggccacacat 174 |
| OY | 121 ctgtctgaatlygagataattaacatcactagaacaacagacaagatgacaataatgtcttaa 180 |
| Db | 175 cgtcgtgaatcgagataattaacatcactagaacaacagacaagatgacaataatgtcttaa 234 |
| OY | 181 gtatgagaatgttttttgacacattccagccctttaatatccacacacagaaagcaac 240 |
| Db | 235 gtatgagaatgttttttgacacattccagccctttaatatccacacacagaaagcaac 294 |
| OY | 241 aaaagaagcacagagatccctctggagaaaatgccggccgcacactcttggtcatcatga 300 |
| Db | 295 aaaagaagcacagagatccctctggagaaaatgccggccgcacactcttggtcatcatga 354 |
| OY | 301 gccctc 305 |
| Db | 355 gccctc 359 |
| | |
| RESULT 11 | |
| XJ37486 | |
| ID XJ37486 standard; cDNA; 597 BP. | |
| XX AC XJ37486; | |
| DT 06-JUL-1999 (first entry) | |
| DE Human secreted protein cDNA fragment containing gene 36. | |
| XX | |
| RW Human; secreted protein; treatment; prevention; protein therapy; AIDS; | |
| RW gene therapy; diagnosis; cancer; tumour; neurodegenerative disorder; | |
| RW developmental abnormality; fetal deficiency; blood disorder; leukemia; | |
| RW immune system disease; autoimmune disease; hepatic disease; lymphoma; | |
| RW renal disease; inflammation; allergy; Alzheimer's disease; schizophrenia; | |
| RW cognitive disorder; prostate disease; skeletal; cardiac; muscle disorder; | |
| RW pulmonary disorder; transplant rejection; osteoclast; osteoporosis; | |
| RW arthritis; malignancy; digestive; endocrine; infection; ss. | |
| XX | |
| OS Homo sapiens. | |
| XX | |
| FN W09918208-A1. | |
| XX | |
| PD 15-APR-1999. | |
| XX | |


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XX HIV-2 genomic DNA sequence.
DE
XX HIV-1; HIV-2; Human immunodeficiency virus; primer/probe set; HIV;
KW detection; nucleic acid amplification; PCR; RT PCR; OH PCR; virus;
KW oligonucleotide hybridisation PCR; ss.
XX
OS Human immunodeficiency virus type 2.
XX
PN WO9558086-A2.
XX
PD 23-DEC-1998.
XX
PF 04-JUN-1998; 98MO-US11652.
XX
PR 16-JUN-1997; 97US-0876546.
XX
PA (ABB0 ) ABBOT LAB.
PI Abiravaya K, Esping CAC, Gorzowski JJ, Hoehnle RJ;
PI Kroegeer PE, Moore JU;
DR WPI: 1999-095352/08.
XX
XX Nucleic acid primers and probes for detecting HIV 1 and HIV 2 -
PT using known nucleic acid amplification procedures especially
PT oligonucleotide hybridisation PCR
XX
XX Disclosure: Page 27-28; 35pp; English.
XX
CC The invention provides primer/probe sets for detecting HIV-1 or HIV-2
CC that comprise two primers and at least one probe. The primer/probe sets
CC are useful to detect HIV (i.e. HIV-1 and HIV-2, either separately or
CC simultaneously) in biological samples using known nucleic acid
CC amplification procedures e.g. PCR, reverse transcriptase (RT) PCR and
CC especially oligonucleotide hybridisation PCR (OH PCR); HIV-1 and HIV-2
CC are different viruses each with several subtypes due to the highly
CC mutable nature of the virus (attributed to the inefficiency with which
CC it converts its genetic material (RNA) into DNA to allow it to insert its
CC genetic information into the host and/or recombination of viral genomes
CC from different HIV populations). The sequences are designed to detect all
CC known HIV-1 or HIV-2 subtypes by OH PCR and allow detection of target
CC sequences which are DNA or, sequences which are embedded within the HIV
CC genome and are therefore RNA by RT-PCR. Sequences X05232 to X05253
CC represent specifically claimed primer and probe sequences present in
CC eight different primer/probe sets that are used for detecting HIV-1 and
CC HIV-2. The present sequence represents HIV-1 genomic DNA sequence that
CC can be used as a target sequence in the method of the invention.
XX
SQ Sequence 2689 BP; 1066 A; 487 C; 597 G; 539 T; 0 other:

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Query Match 5.0%; Score 35.8; DB 20; Length 2689;
Best Local Similarity 53.8%; Pred No. 0.5;
Matches 70; Conservative 0; Mismatches 60; Indels 0; Gaps 0;
OY 561 ggggacagagtcagatctggccctgctgaactgctgataccaatcatlctta 620
DB 1758 GTGCCATTGCAAGGCTTCAATTCGTCGTGATTTGGTTCCTAATACCTTTA 1699
OY 621 ttctacccctaaagctgtngaataatctgaactaaggtcttntggccacatttc 680
DB 1698 CCCTGCTCTCCCTCATCTATATATATCCCTTCCCTCTTTGACTGCCTATTCG 1639
OY 681 atnaccac 690
DB 1638 AGGATCCATC 1629

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